

OIPE

RAW SEQUENCE LISTING

1 <110> APPLICANT: David M. Knipe

PATENT APPLICATION: US/10/050,673

DATE: 02/07/2002 TIME: 14:38:47

_		m		- m				F										
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3					cNam													
	<120>					ION:	Rep	lica	tion	-Com	pete	nt V	irus	Exp	ress	ing	A	
5				Prot														
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7	<140>	CURI	RENT	APP	LICA	TION	NUM	BER:	10/	050,	673							
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	<150>									/127	, 227						DI	<i>ב</i> וי
11	<151>	PRI	OR F	ILIN	G DA	TE:	1998	-07-	31					- N	17		RE	ン
14	<160>	NUM	BER (OF S	EQ I	D NO	S: 6							 '	V I	_	. 1 . –	_
15	<170>	SOF	(WAR	E: F	astS	EQ f	or W	indo	ws V	ersi	on 3	. 0	1	۱ سیا				
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19	<212>	TYPE: DNA																
20	<213>	ORGA	ORGANISM: herpesvirus															
			FEATURE:															
22	<221>	NAME	NAME/KEY: CDS															
			LOCATION: (1)(4359)															
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26																		
27		Met Glu Thr Lys Pro Lys Thr Ala Thr Thr Ile Lys Val Pro Pro Gly 1 5 10 15																
28			cta	ασа	tac	_	tac	act	cac	aca		cca	tcc	gaa	aac		σασ	96
29													Ser					70
30		110	LCu	Ory	20	VUI	1 y 1	AIG	nra	25	Cys	FIO	Ser	GIU	30	TIC	GIU	
31		ctt	cta	aca		cta	toa	aca	cac		aac	a a t	gcc	a a a		acc	ata	144
32													Ala					144
33		пси	LCu	35	пец	Бец	Det	VIG	40	Ser	GIY	изр	Ala	45	vai	мта	Val	
34		aca	ccc		ata	ata	aaa	ata		ata	a 2 a	300	ggc		a 2 a	000	220	192
35													Gly					192
36		мта	50	rea	vai	Val	СТУ	55	1111	Val	GIU	ser	60	Pne	GIU	нта	ASII	
37		at a		a+ a	ata	~+ ~	~~+		~~~	200		~~~		~~~	~~+			240
38													ctc					240
39		65	Ата	Val	Val	Val	70	ser	Arg	THE	THE	_	Leu	GIY	СТА	Thr		
												75					80	000
40													tcg					288
41		Val	ser	Leu	Lys		Thr	Pro	ser	HIS	_	ser	Ser	ser	vaı	-	Val	
42						85					90					95		225
43													cag					336
44		ьиe	HIS	GIA		Arg	HIS	Leu	Asp		ser	Inr	Gln	Ala		Asn	Leu	
45					100					105					110			
46		_	-		_			*-	-				ggc		_	_		384
47		Thr	Arg	Leu	Cys	GLu	Arg	Ala	Arg	Arg	HIS	Phe	Gly	Phe	Ser	Asp	Tyr	

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48			115					120					125				
49								aaa									432
50	Thr	Pro	Arg	Pro	Gly	Asp	Leu	Lys	His	Glu	Thr	Thr	Gly	Glu	Ala	Leu	
51		130					135					140					
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53	Cys	Glu	Arg	Leu	Gly	Leu	Asp	Pro	Asp	Arg	Ala	Leu	Leu	Tyr	Leu	Val	
54	145					150					155					160	
55	qtt	acc	gag	qqc	ttc	aaq	qaq	gcc	qtq	tgc	atc	aac	aac	acc	ttt	ctg	528
56								Ála									
57					165	-				170					175		
58	cac	cta	gga	aac	tca	gac	aaσ	gta	acc	at.a	aac	aaa	aca	σασ	ata	cac	576
59								Val									
60	mis	Lea	017	180	DCI	nop	2,0	, 41	185	110	011	J = 1		190			
61	cac	2 t 2	000		tat	aaa	tta	cag		tta	ato	cca	σat		agg	caa	624
								Gln									021
62	ALG	TIE		vaı	1 1 1	FIO	Leu	200	Leu	FILE	Mec	FIO	205	THE	DCI	my	
63			195								~~~			~~~	~~~	+	672
64								gcc									0/2
65	Val		Ala	GLu	Pro	Pne		Ala	Asn	HIS	Arg		rre	GTÀ	GIU	ASII	
66		210					215					220					700
67								ttt									720
68	Phe	Thr	Tyr	Pro	Leu		Phe	Phe	Asn	Arg		Leu	Asn	Arg	Leu		
69	225					230					235					240	
70								gcc									768
71	Phe	Glu	Ala	Val	Val	Gly	Pro	Ala	Ala	Val	Ala	Leu	Arg	Cys	Arg	Asn	
72					245					250					255		
73	gtg	gac	gcc	gtg	gcc	cgc	gcg	gcc	gcc	cac	ctg	gcg	ttt	gac	gaa	aac	816
74	Val	Asp	Ala	Val	Ala	Arg	Ala	Ala	Ala	His	Leu	Ala	Phe	Asp	Glu	Asn	
75				260					265					270			
76	cac	gag	qqc	qcc	qcc	ctc	ccc	gcc	gac	att	acg	ttc	acg	gcc	ttc	gaa	864
77								Ala									
78			275					280	-				285				
79	gcc	agc	caq	aat	aaq	acc	ccq	cgg	gat	ggg	cqc	qac	qqc	qqc	qqc	aaq	912
80								Arg									
81		290		- 1	1 -		295		_	-	,	300	-	•	_	-	
82	aac		geg	ggc	aaa	ttc		cag	cac	cta	acc	t.cc	at.c	atσ	gcc	gga	960
83								Gln									
84	305		1114	O I I	OI1	310	01.4		5		315					320	
85		acc	acc	cta	acc		σασ	tct	atc	ata		atα	acc	atc	ttc		1008
86								Ser									1000
87	кър	AIG	AIG	Leu	325		GIU	SCI		330		ricc	niu		335		
							+	~~~				+ ~ ~	~ ~ ~				1056
88								gcg									1050
89	GIU	PIO	PIO		ASP	шe	ser	Ala	_	PIO	Leu	Cys	GIU	350	GIII	кэр	
90				340					345	~	~~~	+	at a		000	acc	1104
91								gcc									1104
92	Thr	АІа		Ala	Arg	Ala	Asn	Ala	val	GTA	АТа	ryr		Ald	arg	AIG	
93			355					360					365		- 4		1150
94	gcg	gga	ctc	gtg	ggg	gcc	atg	gta	ttt	agc	acc	aac	tcg	gcc	ctc	cat	1152
95	Ala		Leu	Val	Gly	Ala		Val	Phe	Ser	Thr		Ser	Ala	Leu	His	
96		370					375					380					

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97	ctc	acc	gag	gtg	gac	gac	gcc	ggt	ccg	gcg	gac	сса	aag	gac	cac	agc	1200
98		Thr	Glu	Val	Asp	Asp	Ala	Gly	Pro	Ala	Asp	Pro	Lys	Asp	His	Ser	
99	385					390					395					400	
100																gcg	1248
101	Lys	Pro	Ser	Phe	Tyr	Arg	Phe	Phe	Leu	Val	Pro	Gly	Thr	His	Val	Ala	
102					405					410					415		
103	gcc	aac	cca	cag	gtg	gac	cgc	gag	gga	cac	gtg	gtg	ccc	ggg	ttc	gag	1296
104	Ala	Asn	Pro	Gln	Val	Asp	Arg	Glu	Gly	His	Val	Val	Pro	Gly	Phe	Glu	
105				420					425					430			
106	ggt	cgg	CCC	acc	gcg	CCC	ctc	gtc	ggc	gga	acc	cag	gaa	ttt	gcc	ggc	1344
107	Gly	Arg	Pro	Thr	Ala	Pro	Leu	Val	Gly	Gly	Thr	Gln	Glu	Phe	Ala	Gly	
108			435					440					445				
109																aag	1392
110	Glu	His	Leu	Ala	Met	Leu	Cys	Gly	Phe	Ser	Pro	Ala	Leu	Leu	Ala	Lys	
111		450					455					460					
112	atg	ctg	ttt	tac	ctg	gag	cgc	tgc	gac	ggc	ggc	gtg	atc	gtc	ggg	cgc	1440
113	Met	Leu	Phe	Tyr	Leu	Glu	Arg	Cys	Asp	Gly	Gly	Val	Ile	Val	Gly	Arg	
114	465					470					475					480	
115														cag			1488
116	Gln	Glu	Met	Asp	Val	Phe	Arg	Tyr	Val	Ala	Asp	Ser	Asn	Gln	Thr	Asp	
117					485					490					495		
118														tgc			1536
119	Val	Pro	Cys	Asn	Leu	Cys	Thr	Phe	Asp	Thr	Arg	His	Ala	Cys	Val	His	
120				500					505					510			
121														gcc			1584
122	Thr	Thr		Met	Arg	Leu	Arg	Ala	Arg	His	Pro	Lys	Phe	Ala	Ser	Ala	
123			515					520					525				
124	gcc	cgc	gga	gcc	atc	ggc	gtc	ttc	ggg	acc	atg	aac	agc	atg	tac	agc	1632
125	Ala	Arg	Gly	Ala	Ile	Gly	Val	Phe	Gly	Thr	Met	Asn	Ser	Met	Tyr	Ser	
126		530					535					540					
127														ctg			1680
128		Cys	Asp	Val	Leu		Asn	Tyr	Ala	Ala	Phe	Ser	Ala	Leu	Lys	Arg	
129	545					550					555					560	
130														acg			1728
131	Ala	Asp	Gly	Ser		Thr	Ala	Arg	Thr	Ile	Met	Gln	Glu	Thr	Tyr	Arg	
132					565					570					575		
133														cag			1776
134	Ala	Ala	Thr		Arg	Val	Met	Ala		Leu	Glu	Thr	Leu	Gln	Tyr	Val	
135				580					585					590			
136														atc			1824
137	Asp	Gln		Val	Pro	Thr	Ala		Gly	Arg	Leu	Glu		Ile	Ile	Thr	
138			595					600					605				
139	aac	cgc	gag	gcc	ctg	cat	acg	gtg	gtg	aac	aac	gtc	agg	cag	gtc	gtg	1872
140	Asn		GLu	Ala	Leu	His		Val	Val	Asn	Asn		Arg	Gln	Val	Val	
141		610					615					620					
142	gac	cgc	gag	gtg	gag	cag	ctg	atg	cgc	aac	ctg	gtg	gag	ggg	agg	aac	1920
143		Arg	GLU	va⊥	GLU		Leu	Met.	Arg	Asn		Val	GLu	Gly	Arg		
144	625					630					635					640	
145	LTC	aag	CCC	cgc	gac	ggt	ctg	ggc	gag	gcc	aac	cac	gcc	atg	tcc	ctg	1968

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146 147	Phe	Lys	Phe	Arg	Asp 645	Gly	Leu	Gly	Glu	Ala 650		His	Ala	Met	Ser 655	Leu	
148	acq	cta	gac	cca		aca	tac	aaa	cca			cta	ctt	cag		ctc	2016
149																Leu	2010
150				660	- 1 -		010	011	665	-		204	200	670	1100	Dea	
151	aaa	caa	сда	tcc	aac	ctc	acc	at.a			gac	cta	acc		age	cag	2064
152																Gln	2001
153	1	,	675					680	-			204	685		501	3111	
154	tgc	cac		ata	ttc	qcc	aaa			atc	σασ	aaa		aac	tt.t.	cgc	2112
155																Arg	
156	- 1 -	690	1				695				014	700	9				
157	aat		ttc	caa	cca	ata		caa	caa	cac	at.a		gac	atq	ttt	aac	2160
158	Asn	Gln	Phe	Gln	Pro	Val	Leu	Ara	Ara	Ara	Val	Met	Asp	Met	Phe	Asn	-200
159	705					710		,		,	715		E			720	
160	aac	qqq	ttt	ctq	tcq	qcc	aaa	acq	cta	acq		aca	ctc	tca	σασ		2208
161													Leu				
162		_			725		1			730					735	1	
163	qcq	get	atc	tqc	qcc	ccc	agc	cta	acq	qcc	aac	caq	acg	acc	CCC	qcc	2256
164													Thr				
165				740					745		-			750			
166	gag	agc	aqc	ttc	qaq	qqc	gac	qtt	qcc	cqc	ata	acc	ctg	aaa	ttt	ccc	2304
167													Leu				
168			755			•	-	760		,			765				
169	aag	gag	ctg	cgc	gtc	aag	agc	cgc	gtg	ttg	ttc	aca	ggc	qcq	agc	qcc	2352
170													Gly				
171	_	770		_		_	775					780	-				
172	aac	gcg	tcc	gag	gcc	gcc	aag	gcg	cgg	gtc	gcc	agc	ctc	cag	agc	gcc	2400
173													Leu				
174	785					790					795					800	
175	tac	cag	aag	ccc	gac	aag	cgc	gtg	gac	atc	ctc	ctc	gga	ccg	ctg	ggc	2448
176	Tyr	Gln	Lys	Pro	Asp	Lys	Arg	Val	Asp	Ile	Leu	Leu	Gly	Pro	Leu	Gly	
177					805					810					815		
178													aac				2496
179	Phe	Leu	Leu	Lys	Gln	Phe	His	Ala	Ala	Ile	Phe	Pro	Asn	Gly	Lys	Pro	
180				820					825					830			
181													acg				2544
182	Pro	Gly	Ser	Asn	Gln	Pro	Asn	Pro	Gln	Trp	Phe	Trp	Thr	Ala	Leu	Gln	
183			835					840					845				
184													gac				2592
185	Arg		Gln	Leu	Pro	Ala		Leu	Leu	Ser	Arg	Glu	Asp	Ile	Glu	Thr	
186		850					855					860					
187													gcg				2640
188		Ala	Phe	Ile	Lys		Phe	Ser	Leu	Asp		Gly	Ala	Ile	Asn		
189	865					870					875					880	
190													atg				2688
191	He	Asn	Leu	Ala		Asn	Asn	Val	Ser		Leu	Ala	Met	Tyr	_	Met	
192					885					890					895		
193													tac				2736
194	Ala	ASN	GIN	тте	ьeu	Arg	Tyr	суѕ	Asp	Hls	Ser	Thr	Tyr	Phe	H	Asn	

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196	mb	. CLC	acy	gee	atc	atc	gcg	ggg	tcc	cgc	cgt	. ccc	ccc	agc	gtg	cag	2784
197	1111	теп			тте	rre	Ala		ser	Arg	Arg	Pro		Ser	Val	Gln	
190	~ ~ ~		915					920					925				
	909	geg	gee	gcg	tgg	tcc	gcg	cag	ggc	ggg	gcg	ggc	ctg	gag	gcc	ggg	2832
200	Ата			Ala	Trp	Ser			Gly	Gly	Ala		Leu	Glu	Ala	Gly	
201		930					935					940					
202	gcc	cgc	gcg	ctg	atg	gac	gcc	gtg	gac	gcg	cat	ccg	ggc	gcg	tgg	acg	2880
203			Ala	Leu	Met		Ala	Val	Asp	Ala		Pro	Gly	Ala	Trp	Thr	
204	945					950					955					960	
205	tcc	atg	ttc	gcc	agc	tgc	aac	ctg	ctg	cgg	CCC	gtc	atg	gcg	gcg	cgc	2928
206	Ser	Met	Phe	Ala	Ser	Cys	Asn	Leu	Leu	Arg	Pro	Val	Met	Ala	Ala	Arg	
207					965					970					975		
208	CCC	atg	gtc	gtg	ttg	ggg	ttg	agc	atc	agc	aaa	tac	tac	ggc	atg	gcc	2976
209	Pro	Met	Val	Val	Leu	Gly	Leu	Ser	Ile	Ser	Lys	Tyr	Tyr	Gly	Met	Ala	
210				980					985					990			
211	ggc	aac	gac	cgt	gtg	ttt	cag	gcc	ggg	aac	tgg	gcc	agc	ctg	atg	ggc	3024
212	Gly	Asn	Asp	Arg	Val	Phe	Gln	Ala	Gly	Asn	Trp	Ala	Ser	Leu	Met	Gly	
213			99					100					1009			-	
214	ggc	aaa	aac	gcg	tgc	ccg	ctc	ctt	att	ttt	gac	cgc	acc	cqc	aaq	ttc	3072
215												Arg					
216		101					101				•	1020		-	-		
217	gtc	ctg	gcc	tgt	ccc	cqq	qcc	ggg	ttt	ata	tac	geg	acc	tca	aac	ct.c	3120
218	Val	Leu	Ala	Cys	Pro	Arg	Ála	Gly	Phe	Val	Cys	Ala	Ala	Ser	Asn	Leu	
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220	ggc	qqc	qqa	qcq	cac	qaa	age	tca	cta	tac	σασ	cag	ctc	саа	aac		3168
221	Gly	G1y	Ğĺv	Álá	His	Ğlu	Ser	Ser	Leu	Cvs	Glu	Gln	Leu	Ara	Glv	Tle	3100
222	-	4	-		1045					1050		0	Lou	*** 9	1055		
223	atc	tcc	qaq	gac	aga	aca	acc	atc	acc			gtg	ttc	ata			3216
224												Val					3210
225				1060					1065					1070		1111	
226	ata	aaa	agc	cta	aga	ccc	cac	acc			cta	cag	atc		_	taa	3264
227												Gln					3204
228		-1-	1075		~-1		9	1080		0411	Lea	OIII	1085		пор	111	
229	cta	aca	ctc	ct.a	σασ	σас	αασ			age	gag	gag			gag	cta	3312
230	Leu	Ala	Leu	Leu	Glu	Asp	Glu	Tvr	Leu	Ser	Glu	Glu	Mot	Mot	Clu	Lou	3312
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232	acc			gee	cta	σασ			aac	ממכ	πaπ	tgg		аса	σac	aca	3360
233	Thr	Ala	Ara	Ala	Len	Glu	Ara	Glv	Agn	Glv	Glu	Trp	Sar	Thr	Acn	ycy Ala	2200
234	1105	5	9	u		1110		GLY	пзп	GIY	1115	_	Set	1111	кър	1120	
235			αаσ	ata				acc	nen	aaa		gtc	200	a aa	at a		2400
236	Δla	T.em	Glu	Val	λla	Hie	Clu	λla	Clu	Ala	Lou	Val	Con	Cla	tou	990	3408
237	1114	Dea	Olu	VUI	1125		Giu	AIG	GIU	1130		vaı	ser	GIII	1135		
238	aac	acc	aaa	a a a			330	+++	~~~			~~~	+	~~~			2456
239	auc Δen	Ala	999	gay	9 L 9 17 = 1	Dha	Acc	Dha	999	yaı Na-	LLL Dha	ggc	Lyc	gag	yac	yac Na-	3456
240	nsu	ита	Этү	1140		rne	ASII				Fue	Gly	cys		_	ASP	
241	220	aca	200			aa.	aa.		1145		~~-			1150			2524
242	Acr	315	acy mb∽	Dro	Dha	990 01	99C	ocg n==	999	gcc al-	ccg	gga	ccg	gca	ttt	gcc	350 4
242	ASII	HIG			File	стХ	отА			ита	rro	Gly			Phe	Ala	
443			1155					1160					1165				

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\10050673.raw
Output Set: N:\CRF3\02072002\J050673.raw

L:517 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (6) SEQUENCE: